



FIG. 1A

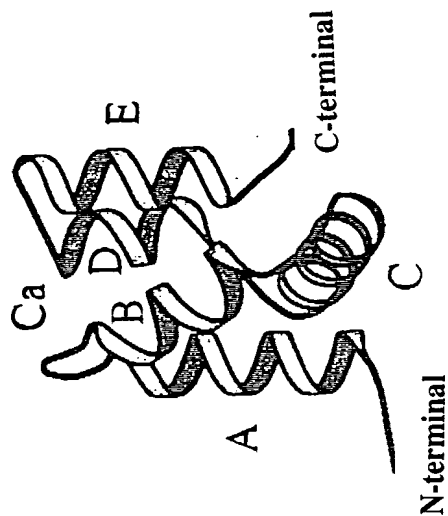
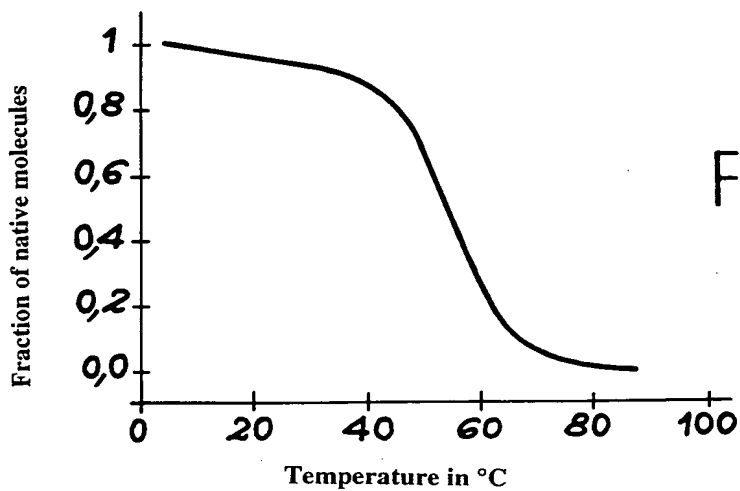
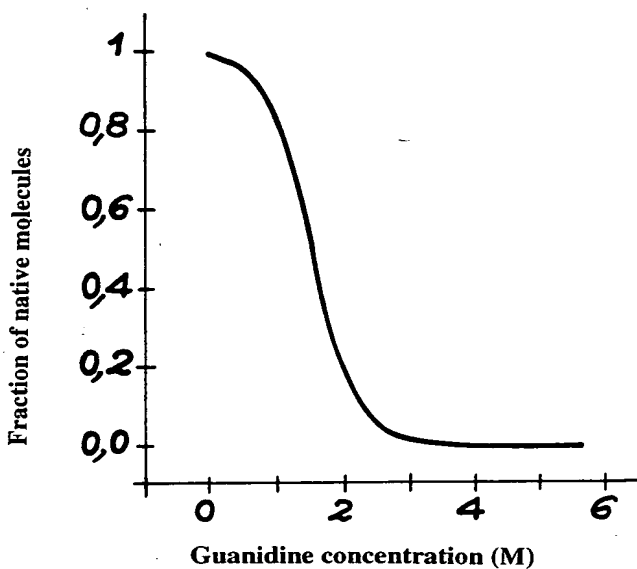
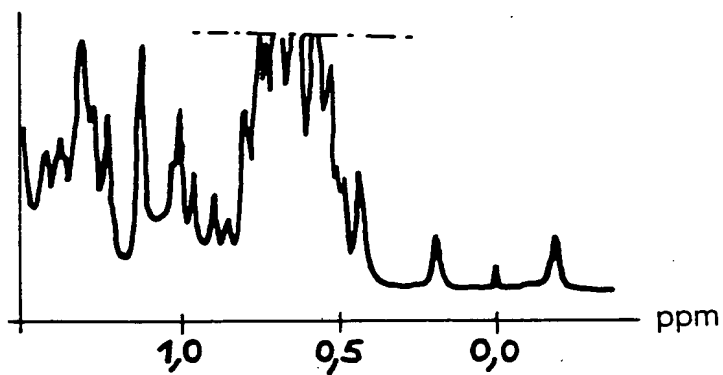


FIG. 1B





Sequence ID No. 1

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Domain 2

Met	Ala	Met	Val	Ser	Glu	Phe	Leu	Lys	Gln	Ala	Trp	Phe	Ile	1	5	10
Glu	Asn	Glu	Glu	Gln	Glu	Tyr	Val	Gln	Thr	Val	Lys	Ser	Ser	15	20	25
Lys	Gly	Gly	Pro	Gly	Ser	Ala	Val	Ser	Pro	Tyr	Pro	Thr	Phe	30	35	40
Asn	Pro	Ser	Ser	Asp	Val	Ala	Ala	Leu	His	Lys	Ala	Ile	Met	45	50	55
Val	Lys	Gly	Val	Asp	Glu	Ala	Thr	Ile	Ile	Asp	Ile	Leu	Thr	60	65	70
Lys	Arg	Asn	Asn	Ala	Gln	Arg	Gln	Gln	Ile	Lys	Ala	Ala	Tyr	75	80	
Leu	Gln	Glu	Thr	Gly	Lys	Pro	Leu	Asp	Glu	Thr	Leu	Lys	Lys	85	90	95
Ala	Leu	Thr	Gly	His	Leu	Glu	Glu	Val	Val	Leu	Ala	Leu	Leu	100	105	110
Lys	Thr	Pro	Ala	Gln	Phe	Asp	Ala	Asp	Glu	Leu	Arg	Ala	Ala	115	120	125
Met	Lys	Gly	Leu	Gly	Thr	Asp	Glu	Asp	Thr	Leu	Ile	Glu	Ile	130	135	140
Leu	Ala	Ser	Arg	Thr	Asn	Lys	Glu	Ile	Arg	Asp	Ile	Asn	Arg	145	150	
Val	Tyr	Arg	Glu	Glu	Leu	Lys	Arg	Asp	Leu	Ala	Lys	Asp	Ile	155	160	165
Thr	Ser	Asp	Thr	Ser	Gly	Asp	Phe	Arg	Asn	Ala	Leu	Leu	Ser	170	175	180
Leu	Ala	Lys	Gly	Asp	Arg	Ser	Glu	Asp	Phe	Gly	Val	Asn	Glu	185	190	200
Asp	Leu	Ala	Asp	Ser	Asp	Ala	Arg	Ala	Leu	Tyr	Glu	Ala	Gly	205	210	215
Glu	Arg	Arg	Lys	Gly	Thr	Asp	Val	Asn	Val	Phe	Asn	Thr	Ile	220	225	
Leu	Thr	Thr	Arg	Ser	Tyr	Pro	Gln	Leu	Arg	Arg	Val	Phe	Gln	230	235	240
Lys	Tyr	Thr	Lys	Tyr	Ser	Lys	His	Asp	Met	Asn	Lys	Val	Leu	245	250	260
Asp	Leu	Glu	Leu	Lys	Gly	Asp	Ile	Glu	Lys	Cys	Leu	Thr	Ala	265	270	275
Ile	Val	Lys	Cys	Ala	Thr	Ser	Lys	Pro	Ala	Phe	Phe	Ala	Glu	280	285	290
Lys	Leu	His	Gln	Ala	Met	Lys	Gly	Val	Gly	Thr	Arg	His	Lys	295	300	
Ala	Leu	Ile	Arg	Ile	Met	Val	Ser	Arg	Ser	Glu	Ile	Asp	Met	305	310	315
Asn	Asp	Ile	Lys	Ala	Phe	Tyr	Gln	Lys	Met	Tyr	Gly	Ile	Ser	320	325	330
Leu	Cys	Gln	Ala	Ile	Leu	Asp	Glu	Thr	Lys	Gly	Asp	Tyr	Glu	335	340	345
Lys	Ile	Leu	Val	Ala	Leu	Cys	Gly	Gly	Asn					350	355	

FIG. 6A: Human annexin I



Sequence ID No. 2

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Domain 1

Met Ala Gln Val Leu Arg Gly Thr Val Thr Asp Phe Pro Gly
1 5 10
Phe Asp Glu Arg Ala Asp Ala Glu Thr Leu Arg Lys Ala Met
15 20 25
Lys Gly Leu Gly Thr Asp Glu Glu Ser Ile Leu Thr Leu Leu
30 35 40
Thr Ser Arg Ser Asn Ala Gln Arg Gln Glu Ile Ser Ala Ala
45 50 55
Phe Lys Thr Leu Phe Gly Arg Asp Leu Leu Asp Asp Leu Lys
60 65 70
Ser Glu Leu Thr Gly Lys Phe Glu Lys Leu Ile Val Ala Leu
75 80
Met Lys Pro Ser Arg Leu Tyr Asp Ala Tyr Glu Leu Lys His
85 90 95
Ala Leu Lys Gly Ala Gly Thr Asn Glu Lys Val Leu Thr Glu
100 105 110
Ile Ile Ala Ser Arg Thr Pro Glu Glu Leu Arg Ala Ile Lys
115 120 125
Gln Val Tyr Glu Glu Glu Tyr Gly Ser Ser Leu Glu Asp Asp
130 135 140
Val Val Gly Asp Thr Ser Gly Tyr Tyr Gln Arg Met Leu Val
145 150
Val Leu Leu Gln Ala Asn Arg Asp Pro Asp Ala Gly Ile Asp
155 160 165
Glu Ala Gln Val Glu Gln Asp Ala Gln Ala Leu Phe Gln Ala
170 175 180
Gly Glu Leu Lys Trp Gly Thr Asp Glu Glu Lys Phe Ile Thr
185 190 195
Ile Phe Gly Thr Arg Ser Val Ser His Leu Arg Lys Val Phe
200 205 210
Asp Lys Tyr Met Thr Ile Ser Gly Phe Gln Ile Glu Glu Thr
215 220
Ile Asp Arg Glu Thr Ser Gly Asn Leu Glu Gln Leu Leu Leu
225 230 235
Ala Val Val Lys Ser Ile Arg Ser Ile Pro Ala Tyr Leu Ala
240 245 250
Glu Thr Leu Tyr Tyr Ala Met Lys Gly Ala Gly Thr Asp Asp
255 260 265
His Thr Leu Ile Arg Val Met Val Ser Arg Ser Glu Ile Asp
270 275 280
Leu Phe Asn Ile Arg Lys Glu Phe Arg Lys Asn Phe Ala Thr
285 290
Ser Leu Tyr Ser Met Ile Lys Gly Asp Thr Ser Gly Asp Tyr
295 300 305
Lys Lys Ala Leu Leu Leu Leu Cys Gly Glu Asp Asp
310 315 320

FIG. 6B Human annexin V



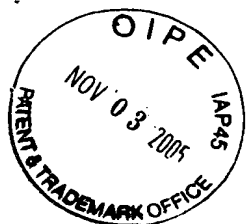
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Sequence ID No. 3

Domain 2

Met	Ala	Ser	Ile	Trp	Val	Gly	His	Arg	Gly	Thr	Val	Arg	Asp
1				5					10				
Tyr	Pro	Asp	Phe	Ser	Pro	Ser	Val	Asp	Ala	Glu	Ala	Ile	Gln
15					20					25			
Lys	Ala	Ile	Arg	Gly	Ile	Gly	Thr	Asp	Glu	Lys	Met	Leu	Ile
	30					35					40		
Ser	Ile	Leu	Thr	Glu	Arg	Ser	Asn	Ala	Gln	Arg	Gln	Leu	Ile
		45					50					55	
Val	Lys	Glu	Tyr	Gln	Ala	Ala	Tyr	Gly	Lys	Glu	Leu	Lys	Asp
			60					65					70
Asp	Leu	Lys	Gly	Asp	Leu	Ser	Gly	His	Phe	Glu	His	Leu	Met
				75					80				
Val	Ala	Leu	Val	Thr	Pro	Pro	Ala	Val	Phe	Asp	Ala	Lys	Gln
85					90					95			
Leu	Lys	Lys	Ser	Met	Lys	Gly	Ala	Gly	Thr	Asn	Glu	Asp	Ala
	100					105					110		
Leu	Ile	Glu	Ile	Leu	Thr	Thr	Arg	Thr	Ser	Arg	Gln	Met	Lys
		115					120					125	
Asp	Ile	Ser	Gln	Ala	Tyr	Tyr	Thr	Val	Tyr	Lys	Lys	Ser	Leu
			130					135					140
Gly	Asp	Asp	Ile	Ser	Ser	Glu	Thr	Ser	Gly	Asp	Phe	Arg	Lys
				145					150				
Ala	Leu	Leu	Thr	Leu	Ala	Asp	Gly	Arg	Arg	Asp	Glu	Ser	Leu
155					160					165			
Lys	Val	Asp	Glu	His	Leu	Ala	Lys	Gln	Asp	Ala	Gln	Ile	Leu
	170					175					180		
Tyr	Lys	Ala	Gly	Glu	Asn	Arg	Trp	Gly	Thr	Asp	Glu	Asp	Lys
		185					190					195	
Phe	Thr	Glu	Ile	Leu	Cys	Leu	Arg	Ser	Phe	Pro	Gln	Leu	Lys
			200					205					210
Leu	Thr	Phe	Asp	Glu	Tyr	Arg	Asn	Ile	Ser	Gln	Lys	Asp	Ile
			215						220				
Val	Asp	Ser	Ile	Lys	Gly	Glu	Leu	Ser	Gly	His	Phe	Glu	Asp
225					230					235			
Leu	Leu	Leu	Ala	Ile	Val	Asn	Cys	Val	Arg	Asn	Thr	Pro	Ala
	240					245					250		
Phe	Leu	Ala	Glu	Arg	Leu	His	Arg	Ala	Leu	Lys	Gly	Ile	Gly
		255					260					270	
Thr	Asp	Glu	Phe	Thr	Leu	Asn	Arg	Ile	Met	Val	Ser	Arg	Ser
			275					280					285
Glu	Ile	Asp	Leu	Leu	Asp	Ile	Arg	Thr	Glu	Phe	Lys	Lys	His
				290					295				
Tyr	Gly	Tyr	Ser	Leu	Tyr	Ser	Ala	Ile	Lys	Ser	Asp	Thr	Ser
300					305					310			
Gly	Asp	Tyr	Glu	Ile	Thr	Leu	Leu	Lys	Ile	Cys	Gly	Gly	Asp
	315					320					325		Asp

FIG. 6C : Human annexin III



Sequence ID No. 4

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OBLON ET AL (703) 413-3000
DOCKET # 205399US0X PCT
INV. Alan SANSON et al.
USSN 09/787,923
Reply to Notice Regarding Drawings
DATED Oct. 17, 2005
REPLACEMENT DRAWINGS

Domain 1

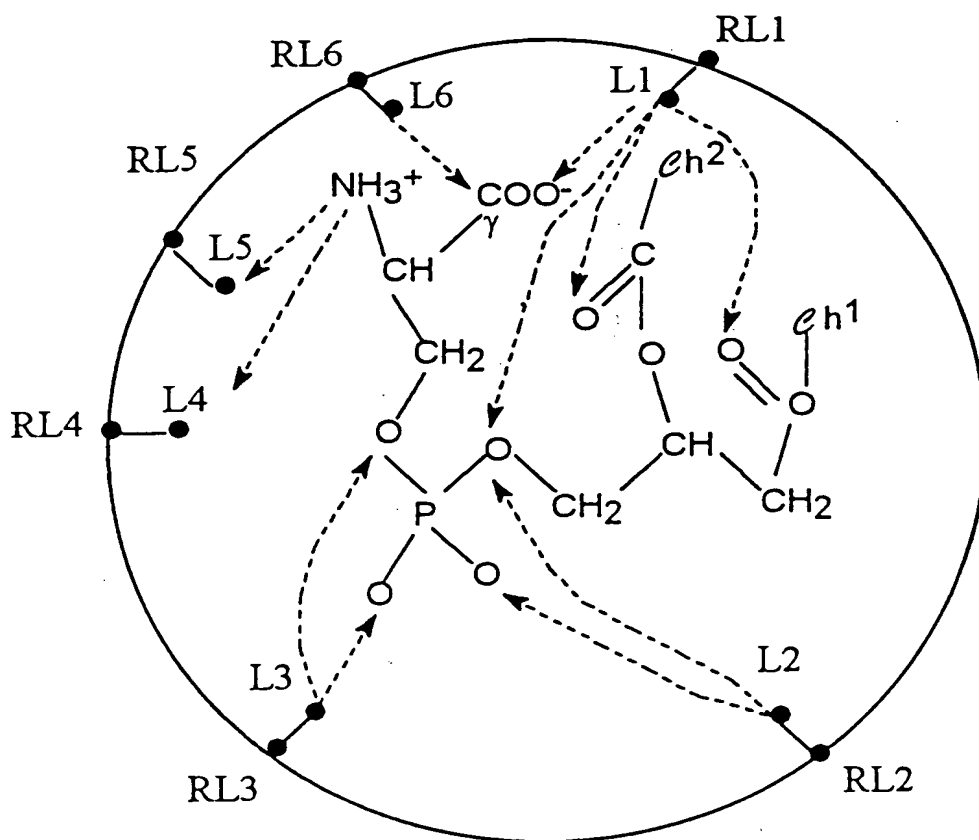
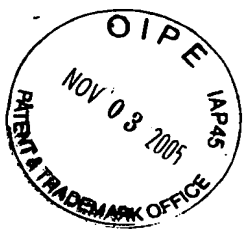
Met	Ala	Thr	Lys	Gly	Gly	Thr	Val	Lys	Ala	Ala	Ser	Gly	Phe
1				5					10				
Asn	Ala	Met	Glu	Asp	Ala	Gln	Thr	Leu	Arg	Lys	Ala	Met	Lys
15				20					25				
Gly	Leu	Gly	Thr	Asp	Glu	Asp	Ala	Ile	Ile	Ser	Val	Leu	Ala
	30			35					40				
Tyr	Arg	Asn	Thr	Ala	Gln	Arg	Gln	Glu	Ile	Arg	Thr	Ala	Tyr
	45			50					55				
Lys	Ser	Thr	Ile	Gly	Arg	Asp	Leu	Ile	Asp	Asp	Leu	Lys	Ser
	60			65					70				
Glu	Leu	Ser	Gly	Asn	Phe	Glu	Gln	Val	Ile	Val	Gly	Met	Met
	75			80									
Thr													
85													

Séquence ID n°5

Domain 2

Pro	Thr	Val	Leu	Tyr	Asp	Val	Gln	Glu	Leu	Gln	Arg	Lys	Gly
86					90					95			
Ala	Met	Lys	Gly	Ala	Gly	Thr	Asp	Glu	Gly	Cys	Leu	Ile	Glu
	100			105						110			
Ile	Leu	Ala	Ser	Arg	Thr	Pro	Glu	Glu	Ile	Arg	Arg	Ile	Asn
	115			120						125			
Gln	Thr	Tyr	Gln	Leu	Gln	Tyr	Gly	Arg	Ser	Leu	Glu	Asp	Asp
	130			135						140			
Ile	Arg	Ser	Asp	Thr	Ser	Phe	Met	Phe	Gln	Arg	Val	Leu	Val
	145			150									
Ser	Leu	Ser	Ala	Gly	Gly	Arg	Asp	Glu	Gly	Asn	Tyr	Leu	Asp
155				160						170			
Asp	Ala	Leu	Val	Arg	Gln	Asp	Ala	Gln	Asp	Leu	Tyr	Glu	Ala
	175			180						185			
Gly	Glu	Lys	Lys	Trp	Gly	Thr	Asp	Glu	Val	Lys	Phe	Leu	Thr
	190			195						200			
Val	Leu	Cys	Ser	Arg	Asn	Arg	Asn	His	Leu	Leu	His	Val	Phe
	205			210						215			
Asp	Glu	Tyr	Lys	Arg	Ile	Ser	Gln	Lys	Asp	Ile	Glu	Gln	Ser
	220			225						230			
Ile	Lys	Ser	Glu	Thr	Ser	Gly	Ser	Phe	Glu	Asp	Ala	Leu	Leu
230				235						240			
Ala	Ile	Val	Lys	Cys	Met	Arg	Asn	Lys	Ser	Ala	Tyr	Phe	Ala
	245			250						255			
Glu	Lys	Leu	Tyr	Lys	Ser	Met	Lys	Gly	Leu	Gly	Thr	Asp	Asp
	260			265						270			
Asn	Thr	Leu	Ile	Arg	Val	Met	Val	Ser	Arg	Ala	Glu	Ile	Asp
	275			280						285			
Met	Leu	Asp	Ile	Arg	Ala	His	Phe	Lys	Arg	Leu	Tyr	Gly	Lys
	290			295						300			
Ser	Leu	Tyr	Ser	Phe	Ile	Lys	Gly	Asp	Thr	Ser	Gly	Asp	Tyr
300				305						310			
Arg	Lys	Val	Leu	Leu	Val	Leu	Cys	Gly	Gly	Asp	Asp		
	315			320						325			

FIG. 6D: Human annexin IV

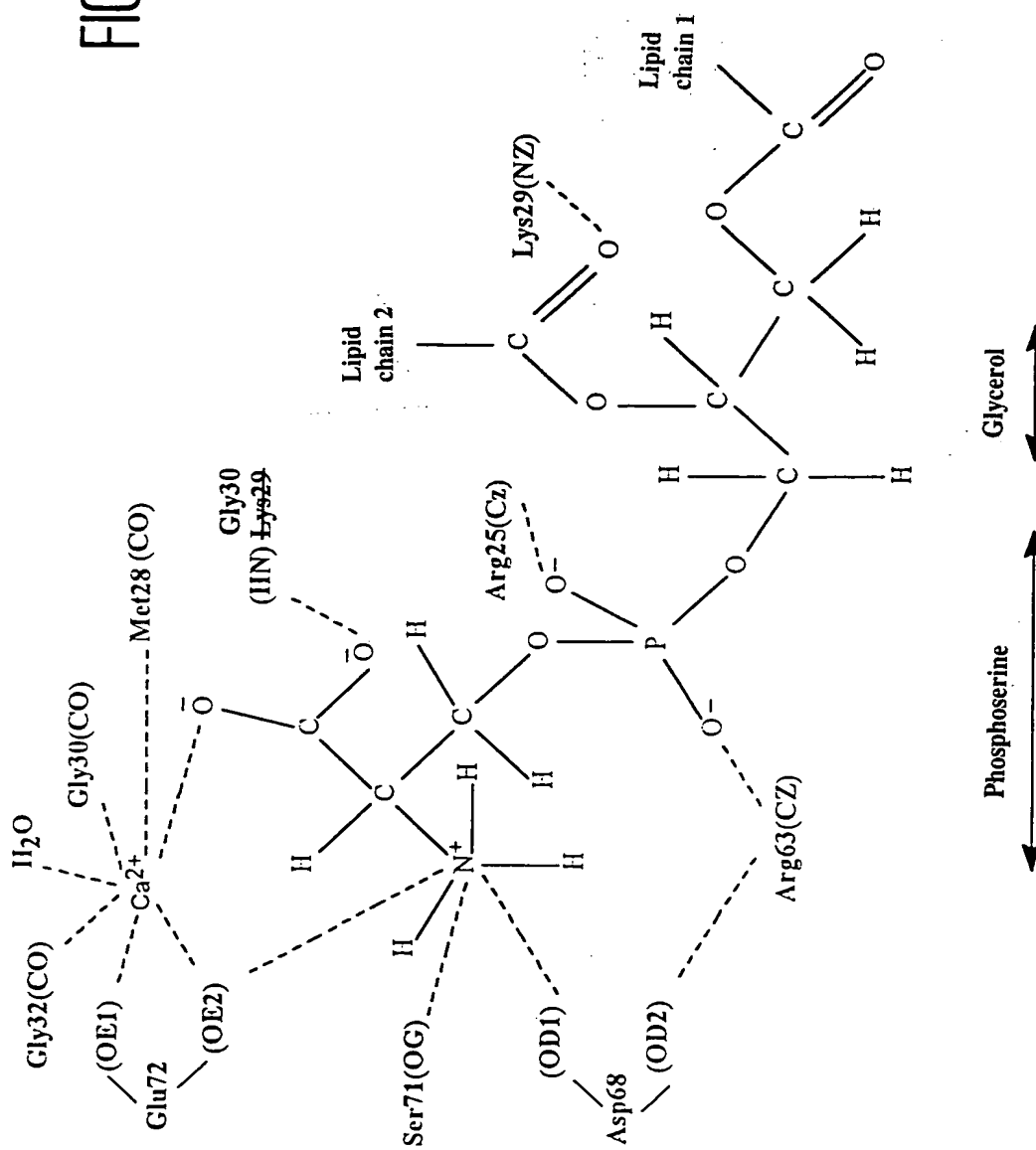


Compound (I) + phosphatidylserine

FIG. 7



FIG. 8





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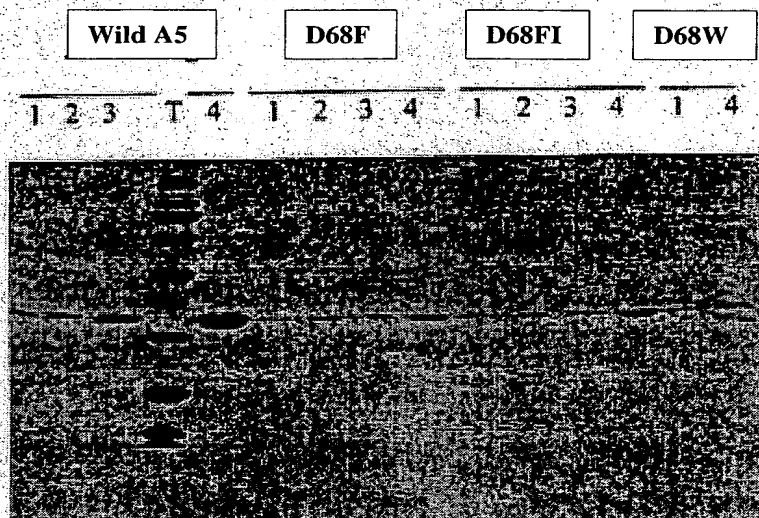


FIG. 9 A

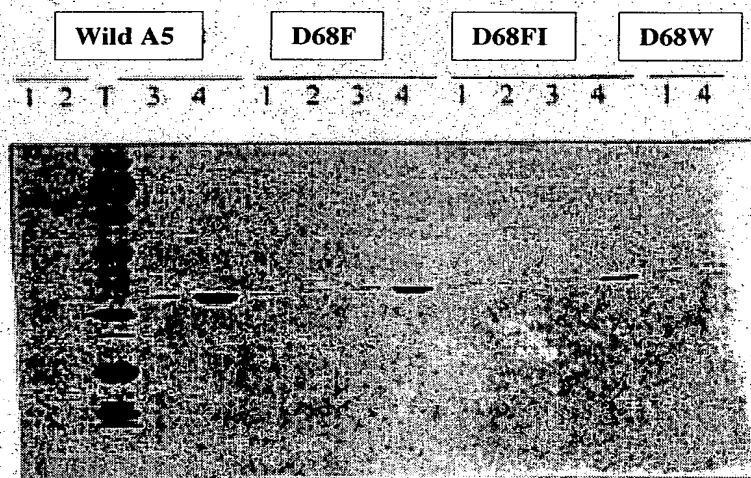


FIG. 9 B

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